

Sequence Listing.ST25
SEQUENCE LISTING

<110> SYRRX, INC.

<120> PROBE, ASSAY AND KITS FOR DETECTING 11 β -HYDROXYSTEROID DEHYDROGENASE AND MODULATORS THEREOF

<130> SYR-HSD-5002

<140> Not Yet Assigned

<141> 2004-03-11

<160> 7

<170> PatentIn version 3.2

<210> 1

<211> 292

<212> PRT

<213> Homo sapiens

<220>

<221> Amino acid sequence for full-length human wild type 11 β -hydroxysteroid dehydrogenase

<222> (1)..(292)

<400> 1

Met Ala Phe Met Lys Lys Tyr Leu Leu Pro Ile Leu Gly Leu Phe Met
1 5 10 15

Ala Tyr Tyr Tyr Tyr Ser Ala Asn Glu Glu Phe Arg Pro Glu Met Leu
20 25 30

Gln Gly Lys Lys Val Ile Val Thr Gly Ala Ser Lys Gly Ile Gly Arg
35 40 45

Glu Met Ala Tyr His Leu Ala Lys Met Gly Ala His Val Val Val Thr
50 55 60

Ala Arg Ser Lys Glu Thr Leu Gln Lys Val Val Ser His Cys Leu Glu
65 70 75 80

Leu Gly Ala Ala Ser Ala His Tyr Ile Ala Gly Thr Met Glu Asp Met
85 90 95

Thr Phe Ala Glu Gln Phe Val Ala Gln Ala Gly Lys Leu Met Gly Gly
100 105 110

Leu Asp Met Leu Ile Leu Asn His Ile Thr Asn Thr Ser Leu Asn Leu
115 120 125

Phe His Asp Asp Ile His His Val Arg Lys Ser Met Glu Val Asn Phe
130 135 140

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Leu Ser Tyr Val Val Leu Thr Val Ala Ala Leu Pro Met Leu Lys Gln
145 150 155 160

Ser Asn Gly Ser Ile Val Val Val Ser Ser Leu Ala Gly Lys Val Ala
165 170 175

Tyr Pro Met Val Ala Ala Tyr Ser Ala Ser Lys Phe Ala Leu Asp Gly
180 185 190

Phe Phe Ser Ser Ile Arg Lys Glu Tyr Ser Val Ser Arg Val Asn Val
195 200 205

Ser Ile Thr Leu Cys Val Leu Gly Leu Ile Asp Thr Glu Thr Ala Met
210 215 220

Lys Ala Val Ser Gly Ile Val His Met Gln Ala Ala Pro Lys Glu Glu
225 230 235 240

Cys Ala Leu Glu Ile Ile Lys Gly Gly Ala Leu Arg Gln Glu Glu Val
245 250 255

Tyr Tyr Asp Ser Ser Leu Trp Thr Thr Leu Leu Ile Arg Asn Pro Cys
260 265 270

Arg Lys Ile Leu Glu Phe Leu Tyr Ser Thr Ser Tyr Asn Met Asp Arg
275 280 285

Phe Ile Asn Lys
290

<210> 2

<211> 286

<212> PRT

<213> Custom

<220>

<221> Amino acid sequence for residues 24-292 of 11 β -hydroxysteroid
dehydrogenase with a N-terminal MKHQHQHQHQHQQQPL tag

<222> (1)..(286)

<400> 2

Met Lys His Gln His Gln His Gln His Gln His Gln Gln Pro
1 5 10 15

Leu Asn Glu Glu Phe Arg Pro Glu Met Leu Gln Gly Lys Lys Val Ile
20 25 30

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Val Thr Gly Ala Ser Lys Gly Ile Gly Arg Glu Met Ala Tyr His Leu
35 40 45

Ala Lys Met Gly Ala His Val Val Val Thr Ala Arg Ser Lys Glu Thr
50 55 60

Leu Gln Lys Val Val Ser His Cys Leu Glu Leu Gly Ala Ala Ser Ala
65 70 75 80

His Tyr Ile Ala Gly Thr Met Glu Asp Met Thr Phe Ala Glu Gln Phe
85 90 95

Val Ala Gln Ala Gly Lys Leu Met Gly Gly Leu Asp Met Leu Ile Leu
100 105 110

Asn His Ile Thr Asn Thr Ser Leu Asn Leu Phe His Asp Asp Ile His
115 120 125

His Val Arg Lys Ser Met Glu Val Asn Phe Leu Ser Tyr Val Val Leu
130 135 140

Thr Val Ala Ala Leu Pro Met Leu Lys Gln Ser Asn Gly Ser Ile Val
145 150 155 160

Val Val Ser Ser Leu Ala Gly Lys Val Ala Tyr Pro Met Val Ala Ala
165 170 175

Tyr Ser Ala Ser Lys Phe Ala Leu Asp Gly Phe Phe Ser Ser Ile Arg
180 185 190

Lys Glu Tyr Ser Val Ser Arg Val Asn Val Ser Ile Thr Leu Cys Val
195 200 205

Leu Gly Leu Ile Asp Thr Glu Thr Ala Met Lys Ala Val Ser Gly Ile
210 215 220

Val His Met Gln Ala Ala Pro Lys Glu Glu Cys Ala Leu Glu Ile Ile
225 230 235 240

Lys Gly Gly Ala Leu Arg Gln Glu Glu Val Tyr Tyr Asp Ser Ser Leu
245 250 255

Trp Thr Thr Leu Leu Ile Arg Asn Pro Cys Arg Lys Ile Leu Glu Phe
260 265 270

Leu Tyr Ser Thr Ser Tyr Asn Met Asp Arg Phe Ile Asn Lys
275 280 285

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<210> 3
<211> 1405
<212> DNA
<213> Custom

<220>
<221> Human cDNA sequence encoding residues 24-292 of
11 β -hydroxysteroid dehydrogenase
<222> (1)..(1405)

<400> 3		
acaattcaga ggctgctgcc tgcttaggag gttgtagaaa gctctgttagg ttctctctgt	60	
gtgtcctaca ggagtcttca ggccagctcc ctgtcgatg gcttttatga aaaaatatct	120	
cctccccatt ctggggctct tcatggccta ctactactat tctgcaaacg aggaattcag	180	
accagagatg ctccaaggaa agaaagtatgat tgtcacaggg gccagcaaag ggatcggaaag	240	
agagatggct tatcatctgg cgaagatggg agcccatgtg gtggtgacag cgaggtcaaa	300	
agaaaactcta cagaaggctgg tatcccactg cctggagctt ggagcagcct cagcacacta	360	
cattgctggc accatgaaag acatgacctt cgcagagcaa tttgttgccc aagcaggaaa	420	
gctcatggga ggactagaca tgctcattct caaccacatc accaacactt ctttgaatct	480	
ttttcatgat gatattcacc atgtgcgaa aagcatgaa gtcaacttcc tcagttacgt	540	
ggtcctgact gtagctgcct tgcccatgct gaagcagagc aatggaagca ttgttgcgt	600	
ctcctctctg gctggaaag tggcttatcc aatggttgct gcctattctg caagcaagtt	660	
tgctttggat gggttttct cctccatcag aaaggaatat tcagtgtcca gggtaatgt	720	
atcaatcact ctctgtgttc ttggcctcat agacacagaa acagccatga aggcagttc	780	
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gggagctctg cgccaaagaag aagtgtatta tgacagctca ctctggacca ctcttctgat	900	
cagaaatcca tgcaggaaga tcctgaaatt tctctactca acgagctata atatggacag	960	
attcataaac aagtaggaac tccctgaggg ctgggcatgc tgagggatt tgggactgtt	1020	
ctgtctcatg tttatctgag ctcttatcta tgaagacatc ttcccagagt gtccccagag	1080	
acatgcaagt catgggtcac acctgacaaa tggaggagt tcctctaaca tttgcaaaat	1140	
ggaaatgtaa taataatgaa tgtcatgcac cgctgcagcc agcagttgta aaattgttag	1200	
taaacatagg tataattacc agatagttat attaaattta tatcttatata ataataat	1260	
gtgatgatta atacaatatt aattataata aaggtcacat aaactttata aattcataac	1320	
tggtagctat aacttgagct tattcaggat ggtttcttta aaaccataaa ctgtacaaat	1380	
gaaatttttc aatatttgtt tctta	1405	

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<211> 24
<212> DNA
<213> Custom

<220>
<221> DNA sequence encoding PCR primer hsd1_24-f
<222> (1)..(24)

<400> 4
aacgaggaat tcagaccaga gatg 24

<210> 5
<211> 24
<212> DNA
<213> Custom

<220>
<221> DNA sequence encoding PCR primer hsd1_292-r
<222> (1)..(24)

<400> 5
ttacttgttt atgaatctgt ccat 24

<210> 6
<211> 23
<212> DNA
<213> Custom

<220>
<221> DNA sequence encoding PCR primer hsdC272sqcf
<222> (1)..(23)

<400> 6
tcagaaatcc atccaggaag atc 23

<210> 7
<211> 23
<212> DNA
<213> Custom

<220>
<221> DNA sequence encoding PCR primer hsdC272Sqcr
<222> (1)..(23)

<400> 7
gatttccctg gatggatttc tga 23